

# Effect of the Enrichment Medium on the Detection and Diversity of *Salmonella* from Porcine Duodenal Content

Emily V. De Busser, Dominiek Maes, Kurt Houf, Jeroen Dewulf, and Lieven De Zutter

## Abstract

This study assesses the effect of the enrichment medium used on the isolation of *Salmonella* from the duodenal content of naturally infected slaughter pigs. At six slaughterhouses, the duodenum was collected from 458 randomly chosen pigs and examined in the laboratory. Three semi-solid enrichment media (modified semi-solid Rappaport-Vassiliadis medium [MSRV], diagnostic semi-solid *Salmonella* medium [DIASALM], and Simple Method *Salmonella* [SMS] agar) and three enrichment broths (Rappaport-Vassiliadis, Rappaport Vassiliadis Broth with Soya [RVS], and Muller Kauffmann Tetrathionate novobiocin broth [MKTTn]) were evaluated. If a migration zone was present on the semi-solid media, a loopful was taken both near the inoculation drop and at the edge of the migration zone and streaked on a Xylose Lysine Desoxycholate (XLD) agar plate. Each enrichment broth was streaked on XLD, and three presumptive colonies were further examined. Detection rate was calculated, and isolates were, after serotyping, genotyped by performing pulsed-field gel electrophoresis (PFGE). The overall frequency of *Salmonella* isolated in at least one of the six different media was 15.5% (71/458). No significant differences in relative sensitivity were obtained within semi-solid media and within liquid media. Semi-solid media showed a significant higher relative sensitivity than the one obtained with liquid media. A relative sensitivity higher than 83.1%, namely of 94.4%, could only be obtained by combining three different enrichment media (MSRV or DIASALM+RVS+MKTTn). In 13.4% of the positive pigs, more than one serotype was found within the duodenum of one pig. In 12.9% of the duodenal contents, different genotypes were found within the same serotype. Differences in serotypes and genotypes were found predominantly within the same enrichment medium. In conclusion, to obtain the highest *Salmonella* detection rate in naturally contaminated pig samples, MSRV should be used as enrichment medium. However, to obtain a realistic picture of the sero- and genotypes present, different samples per enrichment medium and different enrichment media should be tested.

## Introduction

EUROPEAN UNION (EU) DIRECTIVES foresee reduction targets for *Salmonella* in food and animal populations as part of the overall EU strategy to reduce foodborne diseases in humans (Regulation [EC] No 2160/2003). In this context, several surveys obtaining reliable and comparable data on the *Salmonella* prevalence in pigs in EU Member States have already been carried out (Hald *et al.*, 2003; EFSA, 2008, 2010a). Further, national monitoring programs have been established in which bacteriological isolation forms an important part. To obtain comparable results from different countries, it is essential that *Salmonella* isolation procedures are performed accurately and in a standardized manner.

Different media and culture methods are available for the isolation of *Salmonella*. The detection of *Salmonella* spp. in

animal feces and in samples from primary production is described in ISO 6579:2002/Amd 1 2007, in which the selective enrichment medium used is the modified semi-solid Rappaport-Vassiliadis (MSRV), developed for the detection of motile *Salmonella* (EFSA, 2010b) and in which subculture is carried out from the migrated culture, with the inoculum taken from the edge of the visual growth zone. Non-motile *Salmonella* are therefore not isolated. As only *Salmonella* Gallinarum and *Salmonella* Pullorum are not motile (May and Goodner, 1927) and are host specific (poultry), the use of MSRV should not lead to false negative results in *Salmonella* isolation from porcine samples. However, not all salmonellae have the same capacity to migrate on the medium and some may have evolved into non-motile variants (Grimont *et al.*, 2000).

Previous research has shown that the semi-solid media diagnostic semisolid *Salmonella* medium (DIASALM) and

MSRV are not suitable for all *Salmonella* serotypes (O'Donoghue and Winn, 1993; Read *et al.*, 1994). A combination of one of these with the liquid medium Rappaport-Vassiliadis (RV) leads to a higher detection rate of *Salmonella* (Voogt *et al.*, 2001). A further complication is that multiple serotypes can be present in samples from naturally infected animals (Funk *et al.*, 2000; O'Carroll *et al.*, 1999), raising the question whether all of these have an equal chance of being detected (Singer *et al.*, 2009). Probably a particular serotype grows over the others during incubation. This selection pressure has already been described by Harvey and Price's (1967) demonstrating that different *Salmonella* serotypes had different growth characteristics in the same selective enrichment broths. However, Singer *et al.* (2009) suggest that the factors influencing this *in vitro* variability are not solely due to growth competition among *Salmonella* serotypes, as inconsistent results were also found in a fecal experiment tube containing a single *Salmonella* strain.

Further questions involve the number of samples which should be taken and how many suspected colonies derived from the enrichment medium should be tested. These often depend on the objective of the study (e.g., to indicate the samples as *Salmonella* positive or negative, to study the epidemiology of *Salmonella* or to investigate the origin of contaminated food involved in an outbreak). As in the latter two, identifying the sero- and genotypes present is of major importance.

In this study, the impact of six different enrichment media on the detection rate and diversity of *Salmonella* from duodenal content of slaughter pigs is examined.

## Materials and Methods

### Samples

The study was conducted from December 2006 to August 2007. A total of 458 pigs were randomly selected at six different slaughterhouses (A–F). In total, 22 slaughterhouse visits were performed (2–10 visits per slaughterhouse; 10–28 pigs per visit) and 56 different slaughter batches were included. (A slaughter batch contains pigs originating from the same herd.) During evisceration, stomach-gut packages were collected, the duodenum was ligated, taken out, transferred into a sterile bag, and transported to the laboratory.

### *Salmonella* isolation

Upon arrival in the laboratory, the samples were immediately processed for *Salmonella* isolation. Each duodenum was immersed in 95% ethanol and dried in air before being cut open with sterile utensils. Ten grams of duodenal content was diluted 1/10 with buffered peptone water (BPW; Bio-Rad, Marnes-La-Coquette, France), homogenized in a stomacher blender and incubated at 37°C. After 16–20 h, 0.1 mL of the BPW broth was added to 10 mL of RV (Oxoid Ltd., Hampshire, UK) and 10 mL of Rappaport Vassiliadis broth with Soya (RVS; Bio-Rad), spotted on DIASALM (Lab M Ltd., Topley House, Lancashire, UK) and Simple Method *Salmonella* (SMS; AES Chemunex, Bruz Cedex, France) agar, and dispersed in three drops on MRSV (Lab M Ltd. Topley House). Subsequently, 1 mL of the BPW culture was also added to 10 mL of Muller Kauffmann Tetrathionate novo-

biocin broth (MKTTn; Oxoid Ltd.). After incubation of the enrichment media for 24 h at 42°C (but 37°C for MKTTn), the DIASALM, SMS, and MSRV plates were examined for the presence of typical migration zones, and a loopful of the migration zone near the inoculation drop and also from the edge of the migration zone was streaked on a Xylose Lysine Desoxycholate (XLD; Bio-Rad) agar plate. A loopful from each RV, RVS, and MKTTn enrichment broth was also streaked on a XLD agar plate. After incubation for 24 h at 37°C, all XLD plates were examined for the presence of typical colonies, ranging from pink colonies with large, glossy black centers to almost completely black ones. From the semi-solid agars and the enrichment broths, one and three typical colonies, respectively, per XLD plate were selected for identification. In this way, a maximum of 15 colonies per duodenal sample was obtained. Collected isolates were biochemically tested using triple sugar iron, indol, and lysine.

### Evaluation of strain motility

In nine pigs, *Salmonella* could only be isolated after enrichment in one or more broths. The isolates ( $n=18$ ) obtained from these pigs were cultured in Trypto-Casein-Soy Broth (TSB) (Bio-Rad) for 24 h in 37°C. After overnight incubation, two dilutions ( $10^6$  and  $10^3$  cfu/mL) were made in 0.1% peptone water. Of these dilutions, 0.1 mL was spotted on MSRV in three drops and incubated for 24 h at 42°C. The migration capacity of the strains was evaluated by checking the presence of a migration zone on the MSRV agar after incubation. As a reference, 31 isolates obtained from the enrichment broths, but originating from a duodenal sample that tested positive on all enrichment media, were included and submitted to the above mentioned culture method.

### *Salmonella* serotyping

To limit the number of strains that had to be serotyped, all *Salmonella* isolates were subjected to an enterobacterial repetitive intergenic consensus (ERIC) polymerase chain reaction (PCR) (Rasschaert *et al.*, 2005). At least two isolates per cluster were then selected and serotyped by the Belgian reference laboratory for *Salmonella* (Veterinary and

TABLE 1. RELATIVE SENSITIVITIES OF ENRICHMENT MEDIA USED FOR *SALMONELLA* ISOLATION FROM DUODENAL CONTENT OF SLAUGHTER PIGS (71 ITEMS POSITIVE IN AT LEAST ONE METHOD)

Enrichment medium	Number of positive pigs	Relative sensitivity (%)
MSRV	59	83.1
DIASALM	58	81.7
SMS	58	81.7
RV	40	56.3
RVS	39	54.9
MKTTn	34	47.9
All media	71	100

MSRV, modified semi-solid Rappaport-Vassiliadis medium; DIASALM, diagnostic semi-solid *Salmonella* medium; SMS, Simple Method *Salmonella*; RV, Rappaport Vassiliadis; RVS, Rappaport Vassiliadis broth with Soya; MKTTn, Muller Kauffmann Tetrathionate novobiocin broth.

TABLE 2. RESULTS OF McNEMAR TEST AND LEVEL OF AGREEMENT (KAPPA VALUES) FOR DIFFERENT ENRICHMENT MEDIA

		$\kappa$ -Values					
		MSRV	DIASALM	SMS	RV	RVS	MKTTn
p-value McNemar test	MSRV		0.95	0.95	0.74	0.68	0.61
	DIASALM	1.00		0.94	0.75	0.67	0.62
	SMS	1.00	1.00		0.73	0.69	0.62
	RV	<0.01	<0.01	<0.01		0.74	0.65
	RVS	<0.01	<0.01	<0.01	1.00		0.63
	MKTTn	<0.01	<0.01	<0.01	0.31	0.42	

MSRV, modified semi-solid Rappaport-Vassiliadis medium; DIASALM, diagnostic semi-solid *Salmonella* medium; SMS, Simple Method *Salmonella*; RV, Rappaport Vassiliadis; RVS, Rappaport Vassiliadis broth with Soya; MKTTn, Muller Kauffmann Tetrathionate novobiocin broth.

Agrochemical Research Centre, Ukkel, Belgium) using the Kaufmann-White scheme (Popoff and Le Minor, 1992).

### Salmonella genotyping

When more than one isolate with the same serotype was present in a duodenal sample, characterization on strain level was performed by pulsed-field gel electrophoresis (PFGE). The PulseNet protocol (Ribot *et al.*, 2006) was used with the following modification. The running condition was 6 V/cm at 14°C in 0.5× Tris-Borate-EDTA buffer for 20 h with a ramping time from 2.2 to 54.2 seconds. Profiles were obtained by GelCompar II (3.5) (Applied Maths, Sint-Martens-Latem, Belgium) using the Dice coefficient with 2.1% position tolerance, and clusters were generated through the unweighted pair group method using arithmetic averages algorithm (UPGMA). A PFGE genotype was assigned on the basis of a difference in at least one band in the *Xba*I fingerprint. Genotypes within serotypes are indicated below by the capital of the name of the serotype followed by a number (e.g., *Salmonella* Typhimurium genotype 1 is indicated as T<sub>1</sub>).

### Statistical analysis

The relative sensitivity of each culture method was calculated as the number of positive duodenal samples detected by

that method, divided by the total number of samples that tested positive by at least one of the six methods (gold standard).

Possible differences between the proportions of positive pigs obtained with each culture method were analyzed using McNemar tests (SPSS version 19). The level of statistical significance used was  $p < 0.05$ .

To detect the agreement between the results of the culture methods used, kappa values were calculated (SPSS 19). Interpretation of the kappa values was made according to Landis and Koch (1977):  $\leq 0$ , poor agreement; 0.01–0.20, slight agreement; 0.21–0.40, fair agreement; 0.41–0.60, moderate agreement; 0.61–0.80, substantial agreement; 0.81–1.00, almost perfect agreement.

### Results

In 15.5% (71/458) of the samples, *Salmonella* was isolated by at least one of the media. The relative sensitivity of each medium is shown in Table 1. No significant differences were found between the proportion of positive samples obtained by the different semi-solid media or by the different liquid media. However, the proportion of positive samples obtained by the semi-solid media significantly ( $p < 0.05$ ) differed from those obtained by the liquid media (Table 2).

TABLE 3. NUMBER OF PIGS (WITH PERCENTAGES) FOR SPECIFIC *SALMONELLA* SEROTYPES USING DIFFERENT ENRICHMENT MEDIA

Serotype	MSRV	DIASALM	SMS	RV	RVS	MKTTn	Total positive pigs
Total positive pigs	59	58	58	40	39	34	71
<i>Salmonella</i> Typhimurium	41 (69.5%)	41 (70.7%)	38 (65.5%)	25 (62.5%)	19 (48.7%)	21 (61.8%)	45
<i>Salmonella</i> Derby	12 (20.3%)	12 (20.7%)	13 (22.4%)	7 (17.5%)	10 (25.6%)	7 (20.6%)	15
<i>Salmonella</i> Anatum	3 (5.1%)	3 (5.2%)	3 (5.2%)	3 (7.5%)	3 (7.7%)	3 (8.8%)	3
<i>Salmonella</i> Infantis	1 (1.7%)	1 (1.7%)	1 (1.7%)	1 (2.5%)	1 (2.6%)	1 (2.9%)	1
<i>Salmonella</i> Brandenburg	2 (3.4%)	1 (1.7%)	1 (1.7%)	1 (2.5%)	1 (2.6%)	0	2
<i>Salmonella</i> Rissen	3 (5.1%)	1 (1.7%)	2 (3.4%)	1 (2.5%)	1 (2.6%)	0	4
<i>Salmonella</i> O4:i:-	1 (1.7%)	0	0	1 (2.5%)	1 (2.6%)	2 (5.9%)	3
<i>Salmonella</i> Ohio	0	1 (1.7%)	0	1 (2.5%)	1 (2.6%)	0	1
<i>Salmonella</i> Arizonae	0	0	0	0	0	1 (2.9%)	1
Auto-agglutinated	0	0	0	1 (2.5%)	4 (10.2%)	0	4
Total number of identified serotypes/medium	7	7	6	8	8	6	

MSRV, modified semi-solid Rappaport-Vassiliadis medium; DIASALM, diagnostic semi-solid *Salmonella* medium; SMS, Simple Method *Salmonella*; RV, Rappaport Vassiliadis; RVS, Rappaport Vassiliadis broth with Soya; MKTTn, Muller Kauffmann Tetrathionate novobiocin broth.

Calculation of the kappa value reveals an almost perfect agreement between MSRV and DIASALM ( $\kappa=0.95$ ), between MSRV and SMS ( $\kappa=0.95$ ), and between DIASALM and SMS ( $\kappa=0.94$ ). Between all the other media, a substantial ( $\kappa=0.61-0.80$ ) agreement was found (Table 2).

The combined use of MSRV with RVS, MSRV with MKTTn, or DIASALM with RVS increased the detection rate to 63 (88.7%) positive samples. Combining MSRV or DIASALM with RVS and MKTTn led to a rate of 94.4% (67/71 positive duodenal samples).

In total, 113 isolates were collected from the MSRV media, of which 57 were picked close to the inoculation drop and 56 at the edge of the migration zone. On DIASALM and SMS, 51 and 50 isolates, respectively, were picked close to the drop and 55 and 49, respectively, at the edge of the migration zone.

From each slaughterhouse, 4-29 pigs tested positive, and *Salmonella* was isolated in the duodenal content of pigs originating from 28 (50%) slaughter batches. For each batch, an average of 6.5 pigs were sampled, with 2.5 of these (on average) being *Salmonella* positive.

#### Evaluation of strain motility

In four of the nine pigs where *Salmonella* was only isolated after enrichment in one of the broths, serotyping of the isolates ( $n=9$ ) was not possible due to auto-agglutination (Table 3). Six of the nine isolates were able to migrate in MSRV when inoculated at the concentration level of  $10^6$  cfu/mL. When inoculated at the lower concentration level ( $10^3$  cfu/mL), only two did migrate on MSRV. In five pigs, *Salmonella* could only be isolated using RV and MKTTn ( $n=9$ ), and different serotypes were identified (Typhimurium [ $n=3$ ], Arizonae [ $n=3$ ], Derby [ $n=2$ ], and O4:i:- [ $n=1$ ]). Eight of the nine isolates were able to migrate on MSRV when inoculated at the concentration level of  $10^6$  cfu/mL. When inoculated at the lower concentration level ( $10^3$  cfu/mL), this number was declined to three. One of the three *Salmonella* Typhimurium isolates showed no migration capacity when inoculated at both concentration levels.

The 31 isolates selected from duodenal samples where *Salmonella* could be isolated from all the enrichment media showed full migration capacity on MSRV when inoculated at both concentration levels.

#### *Salmonella* serotyping

In total, 595 isolates from the duodenal content of 71 pigs were further serotyped. Table 3 shows the number of pigs for which a particular serotype had been isolated by the different enrichment media used. In general, *Salmonella* Typhimurium was less isolated after enrichment in RVS and *Salmonella* Derby after enrichment in RV. *Salmonella* Anatum was equally detected in all types of enrichment media used, but was in terms of percentage more identified after enrichment in the different broths. *Salmonella* Brandenburg and *Salmonella* Rissen were not isolated after enrichment in MKTTn, while *Salmonella* O4:i:- was not detected after enrichment in DIASALM and SMS. Further, it was not possible to isolate *Salmonella* Ohio from MSRV, SMS, and MKTTn. *Salmonella* Arizonae could only be detected in isolates of MKTTn (Table 3).

No *Salmonella* could be detected from the edge of the migration zone on SMS ( $n=9$ ), MSRV ( $n=3$ ), and DIASALM ( $n=3$ ) in nine animals, while *Salmonella* (Typhimurium, Derby, and Brandenburg) was isolated from cultures taken close

TABLE 4. BREAKDOWN OF NINE PIGS WITH MORE THAN ONE SEROTYPE IN DUODENAL CONTENT ACCORDING TO ENRICHMENT MEDIUM USED

Pig	MSRV		DIASALM		SMS		RV		RVS			MKTTn		
	Close	Edge	Close	Edge	Close	Edge	1	2	3	1	2	1	2	3
7	Derby	Derby	Ohio	Derby	Derby		Ohio			Ohio				
15	Brand	Derby	Derby	Derby	Derby		Derby			Derby		Derby	Derby	
57	Derby	Typhim	Typhim	Typhim	Typhim		Rissen			Derby		O4:i:-	O4:i:-	Derby
278	Typhim	Typhim	Typhim	Typhim	Typhim		Typhim			Typhim		Typhim	Typhim	Typhim
280	O4:i:-	Typhim	Typhim	Typhim	Typhim		O4:i:-			O4:i:-		Typhim	Typhim	Typhim
282	Typhim	Typhim	Typhim	Typhim	Typhim		Typhim			Typhim		Typhim	Typhim	Derby
323	Rissen	Typhim	Typhim	Typhim	Typhim									
325	Typhim	Rissen	Typhim	Rissen	Rissen					Rissen				
327	Typhim	Rissen	Typhim	Typhim	Typhim									

Close, at start of migration zone; edge, at edge of migration zone; Typhim., Typhimurium; Brand., Brandenburg; Rissen, Rappaport-Vassiliadis medium; DIASALM, diagnostic semi-solid *Salmonella* medium; SMS, Simple Method *Salmonella*; RV, Rappaport Vassiliadis; RVS, Rappaport Vassiliadis broth with Soya; MKTTn, Muller Kauffmann Tetrathionate novobiocin broth.



to the inoculation drop. In 15 pigs, *Salmonella* (Typhimurium, Derby, and Rissen) could be obtained from the edge of the migration zone, but not close to the inoculation drop (SMS [ $n=8$ ], DIASALM [ $n=7$ ], and MSRV [ $n=2$ ]).

The number of serotypes found in the isolates of a single slaughter batch varied from one to three. In the majority of the positive pigs (86.6%), only one serotype was present, and in the remaining pigs, two serotypes were present (Table 4). Differences in serotypes amongst all the isolates were found between and within the media used.

### Salmonella genotyping

Genotyping was performed on 578 isolates, resulting in 34 different genotypes. For *Salmonella* Typhimurium, 19 genotypes were obtained, for *Salmonella* Derby five, for *Salmonella* O4:i:- three, for *Salmonella* Rissen and *Salmonella* Brandenburg two, and for the remaining serotypes one. In MSRV, 82.3% of the genotypes were identified; in DIASALM 73.5%; in SMS 64.7%; and in the liquid media RV, RVS, and MKTTn, 58.8%, 44.1%, and 50.0%, respectively. Different genotypes (151 isolates) within and/or between the media were found in the duodenal content of 15 (24.2%) pigs. For seven pigs, the difference in *Salmonella* strain could be explained by a difference in serotype, while for the remaining eight pigs, different genotypes were found within the same serotype. The different genotypes were detected between the different media and within the same medium (Table 5).

The number of genotypes found in the duodenal content of pigs per slaughter batch varied from one to six. The number of different genotypes in each pig varied from one to four, with one genotype found in 77.6% of the pigs, two genotypes in 16.4%, three in 4.5%, and four in 1.5%.

### Discussion

The detection rate of *Salmonella* isolated from naturally contaminated pig samples depends on the enrichment medium used. This study showed that, although semi-solid media are most suitable as enrichment media, a combination of different media is necessary to increase the relative sensitivity. Migration capacity can be influenced by the *Salmonella* serotype (auto-agglutinated or not) and depends on the concentration level of the organism in the sample. Isolates originating from liquid media only were in 77.8% of cases able to

migrate on MSRV when inoculated from a high concentration. At a lower concentration ( $10^3$  cfu/mL), migration capacity decreased to 27.8%. If *Salmonella* is present in the pre-enrichment broth (BPW) at low concentrations, it can easily be missed when semi-solid media are used in the isolation protocol (ISO, 2007). In our procedure, the combination of MSRV or DIASALM with either RVS or MKTTn increased the relative sensitivity from 83.1% (MSRV) or 81.7% (DIASALM) up to 88.7%. These results are in accordance with those obtained by Voogt *et al.* (2001), Dam-Deisz *et al.* (2003), and Botteldoorn *et al.* (2003), indicating that combining media (MSRV or DIASALM with RV) yields a higher number of *Salmonella*-positive samples. The highest relative sensitivity (94.4%) in this study was achieved only by combining three different enrichment media (one semi-solid and two liquid media), increasing the labor intensity and therefore the costs involved.

The distribution of the serotypes showed that some serotypes were less or not recovered from certain media. This finding is also reported by Dam-Deisz *et al.* (2003), although the patterns of serotype detection in the different media used by these authors (MSRV, DIASALM, and RV) are not similar to those found in the present study. Future experimental studies with different serotypes in porcine fecal samples will allow us to gain more insight into the behavior of *Salmonella* serotypes in the standard MSRV medium and also in other enrichment media. Singer *et al.* (2009) showed already that the probability of detecting a specific *Salmonella* serotype in a sample depends on its ability to compete in the cultivation media and on the specific mixture of *Salmonella* bacteria present in the sample. However, explanations for this variability within serotypes need to be further examined.

More than one serotype was identified in the duodenal content of 13.4% of the pigs. The isolation of multiple serotypes from individual pigs has previously been reported (Kampelmacher *et al.*, 1962; O'Carroll *et al.*, 1999; Funk *et al.*, 2000) and is important regarding epidemiological studies. Rostagno *et al.* (2005) demonstrated that asynchronous growth curves among serotypes were due to the selective enrichment media used in the *Salmonella* isolation protocol. In our study, differences in serotypes were predominantly found within the same media, demonstrating that it is useful to select more than one colony per sample. For semi-solid media, samples should be taken both close to the inoculation drop and also on the edge of the migration zone. In this way, less motile *Salmonella* serotypes

TABLE 5. BREAKDOWN OF EIGHT PIGS WITH MORE THAN ONE GENOTYPE WITHIN SAME SEROTYPE ACCORDING TO ENRICHMENT MEDIUM USED

SH	Batch	Pig	MSRV		DIASALM		SMS		RV			RVS			MKTTn		
			Close	Edge	Close	Edge	Close	Edge	1	2	3	1	2	3	1	2	3
B	9	139	T <sub>19</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>9</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>9</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>19</sub>	T <sub>19</sub>	
		142	T <sub>9</sub>	T <sub>13</sub>	T <sub>15</sub>	T <sub>15</sub>	T <sub>15</sub>	T <sub>15</sub>	T <sub>15</sub>	T <sub>15</sub>		T <sub>15</sub>	T <sub>15</sub>		T <sub>15</sub>	T <sub>15</sub>	
	15	273	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>16</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>
		280	O4:i:-1	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	O4:i:-1	T <sub>13</sub>	T <sub>13</sub>	O4:i:-3	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>	T <sub>13</sub>
D	16	314	T <sub>14</sub>	T <sub>12</sub>	T <sub>12</sub>	T <sub>12</sub>	T <sub>14</sub>								T <sub>12</sub>		
	22	325	T <sub>6</sub>	R <sub>2</sub>	T <sub>7</sub>	R <sub>1</sub>		R <sub>1</sub>									
	24	353	T <sub>4</sub>	T <sub>4</sub>		T <sub>4</sub>			T <sub>4</sub>	T <sub>2</sub>	T <sub>4</sub>						
F	28	456	T <sub>14</sub>	T <sub>11</sub>	T <sub>11</sub>	T <sub>14</sub>	T <sub>1</sub>	T <sub>1</sub>	T <sub>14</sub>	T <sub>1</sub>	T <sub>1</sub>				T <sub>1</sub>	T <sub>1</sub>	T <sub>1</sub>

SH, slaughterhouse; Batch, slaughter batch; T, Typhimurium; R, Rissen; close, at start of migration zone; edge, on edge of migration zone; MSRV, modified semi-solid Rappaport-Vassiliadis medium; DIASALM, diagnostic semi-solid *Salmonella* medium; SMS, Simple Method *Salmonella*; RV, Rappaport Vassiliadis; RVS, Rappaport Vassiliadis broth with Soya; MKTTn, Muller Kauffmann Tetrathionate novobiocin broth.

can also be detected. An explanation for the decreased motility of some strains on MSRV cannot be given by the results of this study, as all genotypes (except one) which were found at the start of the migration zone but not on the edge migrated in samples belonging to other pigs. It is possible that the salmonellae were overgrown by competitive bacteria present in the duodenal content of the pig, hindering isolation. In total, 34 strains were found, with the highest diversity within the serotype Typhimurium. The diverse nature of *Salmonella* Typhimurium has also been demonstrated by De Busser *et al.* (2011), showing that in 26 *Salmonella* Typhimurium isolates obtained from the duodenal content of naturally infected slaughtered pigs, six different genotypes were identified. The same study showed that in a total of 162 *Salmonella* Typhimurium isolates obtained from slaughterhouse samples, 32 different genotypes could be detected. Although *Salmonella* Typhimurium has often been considered as very clonal (On and Baggesen, 1997; Gebreyes, 2006), our results illustrate that even without additional phenotypic characterization a wide variety of genotypes belonging to *Salmonella* Typhimurium could be obtained. In 13% of the duodenal content of positive pigs, different strains were found within the same serotype. These differences were predominantly seen within the same medium. This finding again emphasizes the fact that examining more parts of the migration zone of semi-solid media or colonies derived from the enrichment broths increases the probability of detecting multiple genotypes.

## Conclusion

This study shows that even by sampling a rather low number of pigs, multiple sero- and genotypes can be detected within a slaughter batch as well as within one pig. Taking into account the variation resulting from the type of enrichment medium used and the number of colonies taken, it is clear that obtaining realistic and valid data is a real challenge. Although increasing the number of colonies analyzed and using multiple enrichment media inherently involve higher costs and more labor, on occasions when detecting the *Salmonella* source is of primary importance, such as in outbreak investigations, it should be seriously considered.

## Disclosure Statement

No competing financial interests exist.

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Address correspondence to:  
Emily V. De Busser, DVM, PhD  
Faculty of Veterinary Medicine  
Ghent University  
Salisburylaan 133  
B-9820 Merelbeke, Belgium

E-mail: Emily.DeBusser@Ugent.be